



MOUSE Fkh^{sf} cDNA SEQUENCE

1	GCTGATCCCC	CTCTAGCAGT	CCACTTCACC	AAGGTGAGCG	AGTGTCCCTG
51	CTCTCCCCCA	CCAGACACAG	CTCTGCTGGC	GAAAGTGGCA	GAGAGGTATT
101	GAGGGTGGGT	GTCAGGAGCC	CACCAGTACA	GCTGGAAACA	CCCAGCCACT
151	CCAGCTCCCG	GCAACTTCTC	CTGACTCTGC	CTTCAGACGA	GACTTGGAAG
201	ACAGTCACAT	CTCAGCAGCT	CCTCTGCCGT	TATCCAGCCT	GCCTCTGACA
251	AGAACCCAAT	GCCCAACCCT	AGGCCAGCCA	AGCCTATGGC	TCCTTCCTTG
301	GCCCTTGGCC	CATCCCCAGG	AGTCTTGCCA	AGCTGGAAGA	CTGCACCCAA
351	GGGCTCAGAA	CTTCTAGGGA	CCAGGGGCTC	TGGGGGACCC	TTCCAAGGTC
401	GGGACCTGCG	AAGTGGGGCC	CACACCTCTT	CTTCCTTGAA	CCCCCTGCCA
451	CCATCCCAGC	TGCAGCTGCC	TACAGTGCCC	CTAGTCATGG	TGGCACCGTC
501	TGGGGCCCGA	CTAGGTCCCT	CACCCCACCT	ACAGGCCCTT	CTCCAGGACA
551	GACCACACTT	CATGCATCAG	CTCTCCACTG	TGGATGCCCA	TGCCCAGACC
601	CCTGTGCTCC	AAGTGCGTCC	ACTGGACAAC	CCAGCCATGA	TCAGCCTCCC
651	ACCACCTTCT	GCTGCCACTG	GGGTCTTCTC	CCTCAAGGCC	CGGCCTGGCC
701	TGCCACCTGG	GATCAATGTG	GCCAGTCTGG	AATGGGTGTC	CAGGGAGCCA
751	GCTCTACTCT	GCACCTTCCC	ACGCTCGGGT	ACACCCAGGA	AAGACAGCAA
801	CCTTTTGGCT	GCACCCCAAG	GATCCTACCC	${\bf ACTGCTGGCA}$	AATGGAGTCT
851	GCAAGTGGCC	TGGTTGTGAG	AAGGTCTTCG	AGGAGCCAGA	AGAGTTTCTC
901	AAGCACTGCC	${\tt AAGCAGATCA}$	TCTCCTGGAT	GAGAAAGGCA	AGGCCCAGTG
951	CCTCCTCCAG	AGAGAAGTGG	TGCAGTCTCT	GGAGCAGCAG	CTGGAGCTGG
1001	AAAAGGAGAA	GCTGGGAGCT	ATGCAGGCCC	ACCTGGCTGG	GAAGATGGCG
1051	CTGGCCAAGG	CTCCATCTGT	GGCCTCAATG	GACAAGAGCT	CTTGCTGCAT
1101	CGTAGCCACC	AGTACTCAGG	GCAGTGTGCT	CCCGGCCTGG	TCTGCTCCTC
1151	GGGAGGCTCC	AGACGGCGGC	${\tt CTGTTTGCAG}$	${\tt TGCGGAGGCA}$	CCTCTGGGGA
1201	AGCCATGGCA	ATAGTTCCTT	CCCAGAGTTC	TTCCACAACA	TGGACTACTT
1251	CAAGTACCAC	AATATGCGAC	CCCCTTTCAC	CTATGCCACC	CTTATCCGAT
1301	GGGCCATCCT	GGAAGCCCCG	GAGAGGCAGA	GGACACTCAA	TGAAATCTAC
1351	CATTGGTTTA	CTCGCATGTT	CGCCTACTTC	AGAAACCACC	CCGCCACCTG
1401	GAAGAATGCC	ATCCGCCACA	ACCTGAGCCT	GCACAAGTGC	TTTGTGCGAG
1451	TGGAGAGCGA	${\tt GAAGGGAGCA}$	GTGTGGACCG	TAGATGAATT	TGAGTTTCGC
1501	AAGAAGAGGA	GCCAACGCCC	CAACAAGTGC	TCCAATCCCT	GCCCTTGACC
1551	TCAAAACCAA	GAAAAGGTGG	GCGGGGGAGG	GGGCCAAAAC	CATGAGACTG
1601	AGGCTGTGGG	GGCAAGGAGG	CAAGTCCTAC	GTGTACCTAT	GGAAACCGGG
1651	CGATGATGTG	${\tt CCTGCTATCA}$	GGGCCTCTGC	TCCCTATCTA	GCTGCCCTCC
1701	TAGATCATAT	CATCTGCCTT	ACAGCTGAGA	GGGGTGCCAA	TCCCAGCCTA
1751	GCCCCTAGTT	CCAACCTAGC	CCCAAGATGA	ACTTTCCAGT	CAAAGAGCCC
1801	TCACAACCAG	CTATACATAT	CTGCCTTGGC	CACTGCCAAG	CAGAAAGATG
1851	ACAGACACCA	TCCTAATATT	TACTCAACCC	AAACCCTAAA	ACATGAAGAG
1901	CCTGCCTTGG	TACATTCGTG	AACTTTCAAA	GTTAGTCATG	CAGTCACACA
1951	TGACTGCAGT	CCTACTGACT	CACACCCCAA	AGCACTCACC	CACAACATCT
2001	GGAACCACGG	GCACTATCAC	ACATAGGTGT	ATATACAGAC	CCTTACACAG
2051	CAACAGCACT	GGAACCTTCA	CAATTACATC	CCCCCAAACC	ACACAGGCAT
2101	AACTGATCAT	ACGCAGCCTC	AAGCAATGCC	CAAAATACAA	GTCAGACACA
2151	GCTTGTCAGA				

Figure 1

MOUSE Fkhsf PROTEIN SEQUENCE

		-			•
1	MPNPRPAKPM	APSLALGPSP	GVLPSWKTAP	KGSELLGTRG	SGGPFQGRDL
51	RSGAHTSSSL	NPLPPSQLQL	PTVPLVMVAP	SGARLGPSPH	LQALLQDRPH
101	FMHQLSTVDA	HAQTPVLQVR	PLDNPAMISL	PPPSAATGVF	SLKARPGLPP
151	GINVASLEWV	SREPALLCTF	PRSGTPRKDS	NLLAAPQGSY	PLLANGVCKW
201	PGCEKVFEEP	EEFLKHCQAD	HLLDEKGKAQ	CLLQREVVQS	LEQQLELEKE
251	KLGAMQAHLA	GKMALAKAPS	VASMDKSSCC	IVATSTQGSV	LPAWSAPREA
301	PDGGLFAVRR	HLWGSHGNSS	FPEFFHNMDY	FKYHNMRPPF	TYATLIRWAI
351	LEAPERQRTL	NEIYHWFTRM	FAYFRNHPAT	WKNAIRHNLS	LHKCFVRVES
401	EKGAVWTVDE	FEFRKKRSQR	PNKCSNPCP*		•

HUMAN FKH^f cDNA Sequence

1	GCACACACTC	ATCGAAAAAA	ATTTGGATTA	TTAGAAGAGA	GAGGTCTGCG
51				CTCGGTATAA	
101				CAGGCTGATC	
151	AGTCCACTTC	ACCAAGCCTG	CCCTTGGACA	AGGACCCGAT	GCCCAACCCC
201	AGGCCTGGCA	AGCCCTCGGC	CCCTTCCTTG	GCCCTTGGCC	CATCCCCAGG
251				AGCCTCAGAC	
301				GAGATCTTCG	
351				CCACCATCGC	
401				CTCCGGGGCA	
451				ACAGGCCACA	
501				ACCCCTGTGC	
551				CACACCACCC	
601				GCCTCCCACC	
651	GTGGCCAGCC	TGGAATGGGT	GTCCAGGGAG	CCGGCACTGC	TCTGCACCTT
701	CCCAAATCCC	AGTGCACCCA	GGAAGGACAG	CACCCTTTCG	GCTGTGCCCC
751				TCTGCAAGTG	
801				CTCAAGCACT	
851	CCATCTTCTG			ATGTCTCCTC	
901	TGGTACAGTC			TGGAGAAGGA	
951	GCCATGCAGG			GCACTGACCA	
1001	TGTGGCATCA	TCCGACAAGG	GCTCCTGCTG	CATCGTAGCT	GCTGGCAGCC
1051	AAGGCCCTGT	CGTCCCAGCC		CCCGGGAGGC	
1101	CTGTTTGCTG	TCCGGAGGCA	CCTGTGGGGT	AGCCATGGAA	ACAGCACATT
1151	CCCAGAGTTC		TGGACTACTT		AACATGCGAC
1201	CCCCTTTCAC		CTCATCCGCT	GGGCCATCCT	GGAGGCTCCA
1251	GAGAAGCAGC		TGAGATCTAC	CACTGGTTCA	
1301	TGCCTTCTTC		CTGCCACCTG	GAAGAACGCC	ATCCGCCACA
1351	ACCTGAGTCT	GCACAAGTGC	TTTGTGCGGG		
1401	GTGTGGACCG	TGGATGAGCT		AAGAAACGGA	GCCAGAGGCC
1451	CAGCAGGTGT		CACCTGGCCC	CTGACCTCAA	GATCAAGGAA
1501	AGGAGGATGG	ACGAACAGGG		TGGGAGGCAG	
1551				CAGGGACCAA	
1601	TTCCACTGTC			TCCCCCCCTG	
1651	CCTCCCCCAT	CATATCCTTT		TGCTCAGAGG	
1701	CTGGCCCCAG			ACACCCCCCA	
1751	GCAGCCAAAC			CACAGAGCCT	
1801	CTCGCACAGA		GCTGGAAAAG	TCACACAGAC	ACACAAAATG
1851	TCACAATCCT	GTCCCTCAC			

Figure 3

HUMAN FKHsf PROTEIN SEQUENCE

1	MPNPRPGKPS	APSLALGPSP	GASPSWRAAP	KASDLLGARG	PGGTFQGRDL
51	RGGAHASSSS	LNPMPPSQLQ	LPTLPLVMVA	PSGARLGPLP	HLQALLQDRP
101	HFMHQLSTVD	${\bf AHARTPVLQV}$	HPLESPAMIS	LTPPTTATGV	FSLKARPGLP
151		VSREPALLCT		_	
201	WPGCEKVFEE	PEDFLKHCQA	DHLLDEKGRA	QCLLQREMVQ	SLEQQLVLEK
251	EKLSAMQAHL	AGKMALTKAS	SVASSDKGSC	CIVAAGSQGP	VVPAWSGPRE
301	APDSLFAVRR	HLWGSHGNST	FPEFLHNMDY	FKFHNMRPPF	TYATLIRWAI
351	-	NEIYHWFTRM			LHKCFVRVES
401	EKGAVWTVDE	LEFREKRSOR	PSRCSNPTPG	P*	•

Transgenic Vector for generation of FKH^{sf} mice

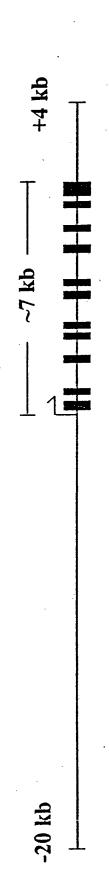


Figure ?

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FKHsf Transgene corrects the defect in scurfy animals

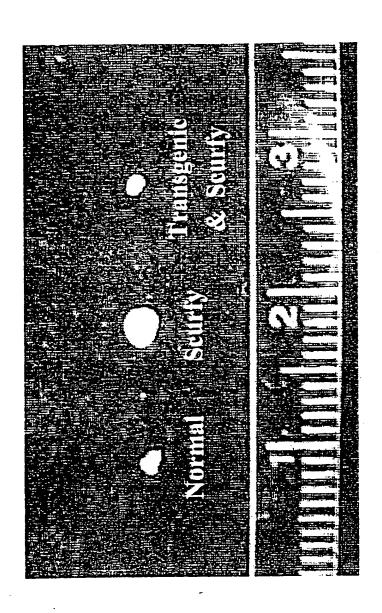


Figure 6

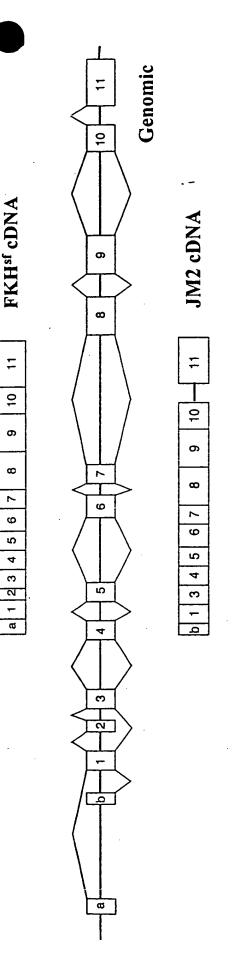
FKHsf tg mice have reduce lymph node cells compared to normal cells

	Mor	Mouse genotype	ype
	Normal	Scurfy	Transgenic
Cell number			
Cells / LN	0.92	1.97	7 0.29
Cells / Thymus	0.76	0.54	4 0.76

FKHsf transgenic mice respond poorly to in vitro stimulation

	Moi	Mouse genotype	ype
	Normal	Scurfy	Transgenic
Proliferation		·	
No stimulation	778	23488	596
Anti-CD3+Anti-CD28	22932	225981	9106

Figure 8



(exons) joined by heavy horizontal lines (introns). Coding exons are numbered 1-11 as determined by sequence analysis of FKH^{sf} cDNA; non-coding 5' exons are labelled a and b. The FKH^{sf}-specific and JM2-specific splicing patterns and Comparison of FKH st and JM2 cDNAs. Exon/intron structure is shown (Genomic) as open rectangles resulting cDNAs are indicated above and below the genomic structure, respectively.

Figure 9

Human FKHsi		Mouse Fkhst
Forkhead	96.4%	·
Mid	82.8%	
ZNE	95.8%	
N-terminal	83.4%	

Human and mouse FKHsf proteins are highly conserved.